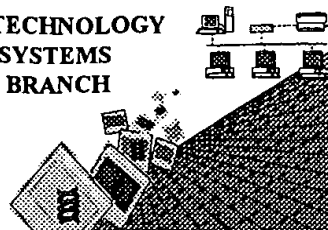


BIOTECHNOLOGY  
SYSTEMS  
BRANCH



**RAW SEQUENCE LISTING**  
**ERROR REPORT**

# 9  
Dm  
3-7c  
**RECEIVED**  
MAR 04 2002  
TECH CENTER 1600/2900

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/922,067B  
Source: OIPE  
Date Processed by STIC: 2/11/2002

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

PATENTIN 2.1 e-mail help: [patin21help@uspto.gov](mailto:patin21help@uspto.gov) or phone 703-306-4119 (R. Wax)

PATENTIN 3.0 e-mail help: [patin3help@uspto.gov](mailto:patin3help@uspto.gov) or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE **CHECKER**  
**VERSION 3.1 PROGRAM**, ACCESSIBLE THROUGH THE U.S. PATENT AND  
TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<<http://www.uspto.gov/ebc/efs/downloads/documents.htm>> , EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: U.S. Patent and Trademark Office, Box Sequence, P.O. Box 2327, Arlington, VA 22202
3. Hand Carry directly to:  
U.S. Patent and Trademark Office, Technology Center 1600, Reception Area, 7<sup>th</sup> Floor, Examiner Name,  
Sequence Information, Crystal Mall One, 1911 South Clark Street, Arlington, VA 22202  
Or  
U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two,  
2011 South Clark Place, Arlington, VA 22202
4. Federal Express, United Parcel Service, or other delivery service to: U.S. Patent and Trademark Office,  
Box Sequence, Room 1B03-Mailroom, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202

Revised 01/29/2002

# Raw Sequence Listing Error Summary

## ERROR DETECTED

## SUGGESTED CORRECTION

SERIAL NUMBER:

09/922,067B

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

- 1      Wrapped Nucleics  
    Wrapped Aminos      The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."
- 2      Invalid Line Length      The rules require that a line not exceed 72 characters in length. This includes white spaces.
- 3      Misaligned Amino  
    Numbering      The numbering under each 5<sup>th</sup> amino acid is misaligned. Do not use tab codes between numbers; use space characters, instead.
- 4      Non-ASCII      The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.
- 5      Variable Length      Sequence(s)        contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.
- 6      PatentIn 2.0  
    "bug"      A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s)       . Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.
- 7      Skipped Sequences  
    (OLD RULES)      Sequence(s)        missing. If intentional, please insert the following lines for each skipped sequence:  
 (2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)  
 (i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading)  
 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)  
 This sequence is intentionally skipped  
 Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.
- 8      Skipped Sequences  
    (NEW RULES)      Sequence(s)        missing. If intentional, please insert the following lines for each skipped sequence.  
 <210> sequence id number  
 <400> sequence id number  
 000
- 9      Use of n's or Xaa's  
    (NEW RULES)      Use of n's and/or Xaa's have been detected in the Sequence Listing.  
 Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present.  
 In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.
- 10      Invalid <213>  
    Response      Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or is Artificial Sequence
- 11      Use of <220>      Sequence(s) 5-7 missing the <220> "Feature" and associated numeric identifiers and responses.  
 Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section.  
 (See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)
- 12      PatentIn 2.0  
    "bug"      Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.
- 13      Misuse of n      n can only be used to represent a single nucleotide in a nucleic acid sequence. N is not used to represent any value not specifically a nucleotide.

AMC/MH - Biotechnology Systems Branch - 08/21/2001



OIFE

RAW SEQUENCE LISTING  
 PATENT APPLICATION: US/09/922,067B

DATE: 02/11/2002  
 TIME: 12:59:47

Input Set : A:\seqlist.txt  
 Output Set: N:\CRF3\02112002\I922067B.raw

Does Not Comply  
 Corrected Diskette Needed

*pp 2-3*

*OK*

4 <110> APPLICANT: MacPhee, Colin Houston  
 5 Tew, David Graham  
 6 Southan, Christopher Donald  
 7 Hickey, Deirdre Mary Bernadette  
 8 Gloger, Israel Simon  
 9 Lawrence, Geoffrey Mark Prouse  
 10 Rice, Simon Quentyn John  
 12 <120> TITLE OF INVENTION: Lipoprotein Associated Phospholipase A2,  
 13 Inhibitors Thereof and Use of the Same in Diagnosis and  
 14 Therapy  
 16 <130> FILE REFERENCE: P30693C4X1C1  
 18 <140> CURRENT APPLICATION NUMBER: US/09/922,067B  
 19 <141> CURRENT FILING DATE: 2001-08-03  
 21 <150> PRIOR APPLICATION NUMBER: 09/193,130  
 22 <151> PRIOR FILING DATE: 2000-11-28  
 24 <150> PRIOR APPLICATION NUMBER: 08/387,858  
 25 <151> PRIOR FILING DATE: 1994-06-24  
 27 <150> PRIOR APPLICATION NUMBER: PCT/GB94/01374  
 28 <151> PRIOR FILING DATE: 1994-06-24  
 30 <150> PRIOR APPLICATION NUMBER: GB 9313144.9  
 31 <151> PRIOR FILING DATE: 1993-06-25  
 33 <160> NUMBER OF SEQ ID NOS: 11  
 35 <170> SOFTWARE: FastSEQ for Windows Version 4.0  
 37 <210> SEQ ID NO: 1  
 38 <211> LENGTH: 37  
 39 <212> TYPE: PRT  
 40 <213> ORGANISM: Homo sapien  
 42 <400> SEQUENCE: 1  
 43 Met Leu Lys Leu Lys Gly Asp Ile Asp Ser Asn Ala Ala Ile Asp Leu  
 44 1 5 10 15  
 45 Ser Asn Lys Ala Ser Leu Ala Phe Leu Gln Lys His Leu Gly Leu His  
 46 20 25 30  
 47 Lys Asp Phe Asp Gln  
 48 35  
 51 <210> SEQ ID NO: 2  
 52 <211> LENGTH: 30  
 53 <212> TYPE: PRT  
 54 <213> ORGANISM: Homo sapien  
 56 <400> SEQUENCE: 2  
 57 Trp Met Phe Pro Leu Gly Asp Glu Val Tyr Ser Arg Ile Pro Gln Pro  
 58 1 5 10 15  
 59 Leu Phe Phe Ile Asn Ser Glu Tyr Phe Gln Tyr Pro Ala Asn  
 60 20 25 30 ~

## RAW SEQUENCE LISTING

DATE: 02/11/2002

PATENT APPLICATION: US/09/922,067B

TIME: 12:59:47

Input Set : A:\seqlist.txt

Output Set: N:\CRF3\02112002\I922067B.raw

63 <210> SEQ ID NO: 3  
 64 <211> LENGTH: 27  
 65 <212> TYPE: PRT  
 66 <213> ORGANISM: Homo sapien  
 68 <400> SEQUENCE: 3  
 69 Gln Tyr Ile Asn Pro Ala Val Met Ile Thr Ile Arg Gly Ser Val His  
 70 1 5 10 15  
 71 Gln Asn Phe Ala Asp Phe Thr Phe Ala Thr Gly  
 72 20 25

75 <210> SEQ ID NO: 4  
 76 <211> LENGTH: 19  
 77 <212> TYPE: PRT  
 78 <213> ORGANISM: Homo sapien  
 80 <400> SEQUENCE: 4  
 81 Trp Leu Met Gly Asn Ile Leu Arg Leu Leu Phe Gly Ser Met Thr Thr  
 82 1 5 10 15  
 83 Pro Ala Asn

87 &lt;210&gt; SEQ ID NO: 5

88 &lt;211&gt; LENGTH: 420

89 &lt;212&gt; TYPE: DNA

90 &lt;213&gt; ORGANISM: Unknown

92 &lt;220&gt; FEATURE:

93 &lt;223&gt; OTHER INFORMATION: Where N can be represented by A, C, T, or G

95 &lt;221&gt; NAME/KEY: misc\_feature

96 &lt;222&gt; LOCATION: 265, 390, 395, 403, 406

97 &lt;223&gt; OTHER INFORMATION: n = A,T,C or G

99 &lt;400&gt; SEQUENCE: 5

100 aaaaaaccta ttttaacct aattgtatct ctctattcct gaagagttct gtaacatgat 60  
 101 gtgttgattg gttgtgttaa tggtgtccc tggataaga ttctcatcat ctccttcaat 120  
 102 caagcagtc cactgatcaa aatctttatg aagtcctaaa tgcttttgta agaattgctaa 180  
 103 tgaagctttg ttgctaagat caatagctgc atttgaatct atgtctccct ttaatttgag 240  
 104 catgtgtcca attattttgc cagtgcaaaa agtgaagtca gcaaaattct ggtggactga 300  
 105 acccctgatt gtaatcatct ttctttcttt atcaggtgag tagcattttt tcatttttat 360  
 106 gatattagca ggatattgga aatattcag gttgntaaaa agngngnggt gagggattct 420

109 &lt;210&gt; SEQ ID NO: 6

110 &lt;211&gt; LENGTH: 379

111 &lt;212&gt; TYPE: DNA

112 &lt;213&gt; ORGANISM: Unknown

114 &lt;220&gt; FEATURE:

115 &lt;223&gt; OTHER INFORMATION: Where N can be represented by A, C, T, or G

117 &lt;221&gt; NAME/KEY: misc\_feature

118 &lt;222&gt; LOCATION: 84

119 &lt;223&gt; OTHER INFORMATION: n = A,T,C or G

121 &lt;400&gt; SEQUENCE: 6

122 tgctaataatc ataaaaatga aaaaatgcta ctcacctgat aaagaaagaa agatgattac 60  
 123 aatcaggggt tcagtcacc aganttttgc tgacttcact tttgcaactg gcaaaataat 120  
 124 tggacacatg ctcaaattaa agggagacat agattcaaatt gtagctattg atcttagcaa 180  
 125 caaagcttca ttagcattct taaaaagca tttaggactt cataaagatt ttgttcagtg 240  
 126 ggactgcttg attgaaggag atgatgagaa tcttattcca ggaccaaca ttaacacaac 300

## RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/922,067B.

DATE: 02/11/2002

TIME: 12:59:47

Input Set : A:\seqlist.txt

Output Set: N:\CRF3\02112002\I922067B.raw

127 caattcaaca catcatgttt acagaacttc ttccaggga taggaggaaa tacaattggg 360  
 128 gtttaaaata ggtttttt 379  
 130 <210> SEQ ID NO: 7  
 131 <211> LENGTH: 279  
 132 <212> TYPE: DNA  
 133 <213> ORGANISM: Unknown *dem 11*  
 135 <220> FEATURE:  
 136 <223> OTHER INFORMATION: Where N can be represented by A, C, T, or G  
 138 <221> NAME/KEY: misc\_feature  
 139 <222> LOCATION: 257  
 140 <223> OTHER INFORMATION: n = A,T,C or G  
 142 <400> SEQUENCE: 7  
 143 gaagaatgca ttagatttaa agtttgatat ggaacaactg aaggactcta ttgataggga 60  
 144 aaaaatagca gtaattggac attcttttgg tggagcaacg gttattcaga ctcttagtga 120  
 145 agatcagaga ttcagatgtg gtattgccct ggatgcatgg atgtttccac tgggtgatga 180  
 146 agtatattcc agaattcctc agccctcttt ttttatcaac tctgaatatt tccaatatcc 240  
 147 *W* tgctaataatc ataaaantgg aaaaatgcta ctcacctgg 279  
 149 <210> SEQ ID NO: 8  
 150 <211> LENGTH: 572  
 151 <212> TYPE: DNA  
 152 <213> ORGANISM: Homo sapien  
 154 <400> SEQUENCE: 8  
 155 aaaatagcag taattggaca ttcttttaggt ggagcaacgg ttattcagac tcttagtgaa 60  
 156 gatcagagat tcagatgtgg tattgccctg gatgcatgga tgtttccact gggtgatgaa 120  
 157 gtatattcca gaattcctca gcccctcttt tttatcaact ctgaatattt ccaatatcct 180  
 158 gctaatatca taaaaatgaa aaaatgctac tcacctgata aagaaagaaa gatgattaca 240  
 159 atcaggggtt cagtccacca gaattttgct gacttcaact ttgcaactgg caaaataatt 300  
 160 ggacacatgc tcaaattaaa gggagacata gattcaaatt tagctattga tcttagcaac 360  
 161 aaagcttcat cagcattctt acaaaagcat ttaggacttc ataaagattt tgatcagtgg 420  
 162 gactgcttga ttgaaggaga tgatgagaat ottattccag ggaccaacat taacacaacc 480  
 163 aatcaacaca tcatgttaca gaactcttca ggaatagaga aatacaatta ggattaaaat 540  
 164 aggtttttta aaaaaaaaaa aaaaaaaact cg 572  
 166 <210> SEQ ID NO: 9  
 167 <211> LENGTH: 1361  
 168 <212> TYPE: DNA  
 169 <213> ORGANISM: Homo sapien  
 171 <220> FEATURE:  
 172 <221> NAME/KEY: CDS  
 173 <222> LOCATION: (38)...(1360)  
 175 <400> SEQUENCE: 9  
 176 ttagagacta agctgaaact gctgctcagc tcccaag atg gtg cca ccc aaa ttg 55  
 177 Met Val Pro Pro Lys Leu  
 178 1 5  
 180 cat gtg ctt ttc tgc ctc tgc ggc tgc ctg gct gtg gtt tat cct ttt 103  
 181 His Val Leu Phe Cys Leu Cys Gly Cys Leu Ala Val Val Tyr Pro Phe  
 182 10 15 20  
 184 gac tgg caa tac ata aat cct gtt gcc cat atg aaa tca tca gca tgg 151  
 185 Asp Trp Gln Tyr Ile Asn Pro Val Ala His Met Lys Ser Ser Ala Trp  
 186 25 30 35

## RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/922,067B

DATE: 02/11/2002

TIME: 12:59:47

Input Set : A:\seqlist.txt

Output Set: N:\CRF3\02112002\I922067B.raw

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188 gtc aac aaa ata caa gta ctg atg gct gct gca agc ttt ggc caa act 199
189 Val Asn Lys Ile Gln Val Leu Met Ala Ala Ala Ser Phe Gly Gln Thr
190      40      45      50
192 aaa atc ccc cgg gga aat ggg cct tat tcc gtt ggt tgt aca gac tta 247
193 Lys Ile Pro Arg Gly Asn Gly Pro Tyr Ser Val Gly Cys Thr Asp Leu
194 55      60      65      70
196 atg ttt gat cac act aat aag ggc acc ttc ttg cgt tta tat tat cca 295
197 Met Phe Asp His Thr Asn Lys Gly Thr Phe Leu Arg Leu Tyr Tyr Pro
198      75      80      85
200 tcc caa gat aat gat cgc ctt gac acc ctt tgg atc cca aat aaa gaa 343
201 Ser Gln Asp Asn Asp Arg Leu Asp Thr Leu Trp Ile Pro Asn Lys Glu
202      90      95      100
204 tat ttt tgg ggt ctt agc aaa ttt ctt gga aca cac tgg ctt atg ggc 391
205 Tyr Phe Trp Gly Leu Ser Lys Phe Leu Gly Thr His Trp Leu Met Gly
206      105      110      115
208 aac att ttg agg tta ctc ttt ggt tca atg aca act cct gca aac tgg 439
209 Asn Ile Leu Arg Leu Leu Phe Gly Ser Met Thr Thr Pro Ala Asn Trp
210      120      125      130
212 aat tcc cct ctg agg cct ggt gaa aaa tat cca ctt gtt gtt ttt tct 487
213 Asn Ser Pro Leu Arg Pro Gly Glu Lys Tyr Pro Leu Val Val Phe Ser
214 135      140      145      150
216 cat ggt ctt ggg gca ttc agg aca ctt tat tct gct att ggc att gac 535
217 His Gly Leu Gly Ala Phe Arg Thr Leu Tyr Ser Ala Ile Gly Ile Asp
218      155      160      165
220 ctg gca tct cat ggg ttt ata gtt gct gct gta gaa cac aga gat aga 583
221 Leu Ala Ser His Gly Phe Ile Val Ala Ala Val Glu His Arg Asp Arg
222      170      175      180
224 tct gca tct gca act tac tat ttc aag gac caa tct gct gca gaa ata 631
225 Ser Ala Ser Ala Thr Tyr Tyr Phe Lys Asp Gln Ser Ala Ala Glu Ile
226      185      190      195
228 ggg gac aag tct tgg ctc tac ctt aga acc ctg aaa caa gag gag gag 679
229 Gly Asp Lys Ser Trp Leu Tyr Leu Arg Thr Leu Lys Gln Glu Glu Glu
230      200      205      210
232 aca cat ata cga aat gag cag gta cgg caa aga gca aaa gaa tgt tcc 727
233 Thr His Ile Arg Asn Glu Gln Val Arg Gln Arg Ala Lys Glu Cys Ser
234 215      220      225      230
236 caa gct ctc agt ctg att ctt gac att gat cat gga aag cca gtg aag 775
237 Gln Ala Leu Ser Leu Ile Leu Asp Ile Asp His Gly Lys Pro Val Lys
238      235      240      245
240 aat gca tta gat tta aag ttt gat atg gaa caa ctg aag gac tct att 823
241 Asn Ala Leu Asp Leu Lys Phe Asp Met Glu Gln Leu Lys Asp Ser Ile
242      250      255      260
244 gat agg gaa aaa ata gca gta att gga cat tct ttt ggt gga gca acg 871
245 Asp Arg Glu Lys Ile Ala Val Ile Gly His Ser Phe Gly Gly Ala Thr
246      265      270      275
248 gtt att cag act ctt agt gaa gat cag aga ttc aga tgt ggt att gcc 919
249 Val Ile Gln Thr Leu Ser Glu Asp Gln Arg Phe Arg Cys Gly Ile Ala
250      280      285      290
252 ctg gat gca tgg atg ttt cca ctg ggt gat gaa gta tat tcc aga att 967

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## RAW SEQUENCE LISTING

DATE: 02/11/2002

PATENT APPLICATION: US/09/922,067B

TIME: 12:59:47

Input Set : A:\seqlist.txt

Output Set: N:\CRF3\02112002\I922067B.raw

```

253 Leu Asp Ala Trp Met Phe Pro Leu Gly Asp Glu Val Tyr Ser Arg Ile
254 295          300          305          310
256 cct cag ccc ctc ttt ttt atc aac tct gaa tat ttc caa tat cct gct 1015
257 Pro Gln Pro Leu Phe Phe Ile Asn Ser Glu Tyr Phe Gln Tyr Pro Ala
258          315          320          325
260 aat atc ata aaa atg aaa aaa tgc tac tca cct gat aaa gaa aga aag 1063
261 Asn Ile Ile Lys Met Lys Lys Cys Tyr Ser Pro Asp Lys Glu Arg Lys
262          330          335          340
264 atg att aca atc agg ggt tca gtc cac cag aat ttt gct gac ttc act 1111
265 Met Ile Thr Ile Arg Gly Ser Val His Gln Asn Phe Ala Asp Phe Thr
266          345          350          355
268 ttt gca act ggc aaa ata att gga cac atg ctc aaa tta aag gga gac 1159
269 Phe Ala Thr Gly Lys Ile Ile Gly His Met Leu Lys Leu Lys Gly Asp
270          360          365          370
272 ata gat tca aat gca gct att gat ctt agc aac aaa gct tca tta gca 1207
273 Ile Asp Ser Asn Ala Ala Ile Asp Leu Ser Asn Lys Ala Ser Leu Ala
274 375          380          385          390
276 ttc tta caa aag cat tta gga ctt cat aaa gat ttt gat cag tgg gac 1255
277 Phe Leu Gln Lys His Leu Gly Leu His Lys Asp Phe Asp Gln Trp Asp
278          395          400          405
280 tgc ttg att gaa gga gat gat gag aat ctt att cca ggg acc aac att 1303
281 Cys Leu Ile Glu Gly Asp Asp Glu Asn Leu Ile Pro Gly Thr Asn Ile
282          410          415          420
284 aac aca acc aat caa cac atc atg tta cag aac tct tca gga ata gag 1351
285 Asn Thr Thr Asn Gln His Ile Met Leu Gln Asn Ser Ser Gly Ile Glu
286          425          430          435
288 aaa tac aat t 1361
289 Lys Tyr Asn
290          440
293 <210> SEQ ID NO: 10
294 <211> LENGTH: 7
295 <212> TYPE: PRT
296 <213> ORGANISM: Homo sapien
298 <400> SEQUENCE: 10
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300 1          5
303 <210> SEQ ID NO: 11
304 <211> LENGTH: 20
305 <212> TYPE: PRT
306 <213> ORGANISM: Homo sapien
308 <400> SEQUENCE: 11
309 Met Ile Thr Ile Arg Gly Ser Val His Gln Asn Phe Ala Asp Phe Thr
310 1          5          10          15
311 Phe Ala Thr Gly
312          20

```

## VERIFICATION SUMMARY

PATENT APPLICATION: US/09/922,067B

DATE: 02/11/2002

TIME: 12:59:48

Input Set : A:\seqlist.txt

Output Set: N:\CRF3\02112002\I922067B.raw

L:18 M:270 C: Current Application Number differs, Wrong Format

L:104 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5

L:106 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5

L:123 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6

L:147 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7